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## CLAIMS

niae PBP2x, characterized in that it consists of a concatenation of the fragments corresponding respectively to the amino acids located between positions 74 to 90, 186 to 199, 218 to 228 and 257-750, with reference to the sequence of the PBP2x protein of the strain R6 (SWISSPROT P14677 or GENBANK 18266817), each one of said fragments being preceded by a peptide fragment of 1 to 7 amino acids.

2. Protein according to Claim 1, characterized in that said peptide fragment comprises amino acids of said Streptococcus pneumoniae PBP2x protein located between positions -1 to -7, relative to the residues at positions 74, 186, 218 and 257, and/or between positions +1 to +7, relative to the residues at positions 90, 199 and 228, as defined in Claim 1.

3.Protein according to Claim 1 or Claim 2, characterized in that said peptide fragment comprises amino acids chosen from alanine (A), serine (S), glycine (G) and threonine (T).

4.Protein according to any one of Claims 1 to 3, characterized in that it is derived from a  $\beta$ -lactam-resistant strain of s. pneumoniae.

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- 5.Protein according to any one of Claims 1 to 3, characterized in that it has the sequence SEQ ID No. 1.
- 6. Protein according to any one of Claims 1 to 5, characterized in that it comprises a substitution of at least one methionine residue with a selenomethionine residue.
  - 7. Protein according to any one of Claims 1 to 6, characterized in that it is associated with a ligand.
  - 8. Protein according to any one of Claims 1 to 7, characterized in that it is in the form of a crystal.
  - 9.Peptide, characterized in that it consists of a fragment of at least 7 amino acids of the mini-PBP2x protein, according to any one of Claims 1 to 6, which peptide includes at least one residue chosen from those located at positions 74, 90, 186, 199, 218, 228 and 257 as defined in Claim 1.
  - 10.Antibodies, characterized in that they are directed against a peptide according to Claim 9.
  - 11. Isolated nucleic acid molecule, characterized in that it is selected from the group consisting of the sequences encoding a mini-PBP2x according to any one of Claims 1 to 6 and the sequences complementary to the preceding sequences, which are sense or antisense.

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- 12. Pair of primers, characterized in that it has the sequence SEQ ID Nos. 2-3.
- Primers, characterized in that they comprise a sequence of approximately 10 to 30 nucleotides 5 corresponding to that located at the junction of the peptide fragments of 1 to 7 amino acids and the fragments of PBP2x as defined in Claim 1.
- 14. Primers according to Claim 13, characterized in that they have a sequence selected from the group consisting of the sequences SEQ ID Nos. 4 to 9. 10
  - 15. Recombinant vector, characterized in that it comprises an insert selected from the group consisting of the nucleic acid molecules encoding a mini-PBP2x according to Claim 11.
- 16. Expression vector according to Claim 15, 15 characterized in that it consists of a prokaryotic vector.
  - 17. Cells transformed with a recombinant vector according to either one of Claims 15 and 16.
- 18.Cells according to Claim 17, characterized 20 in that they are prokaryotic cells.
  - 19. Use of a mini-PBP2x according to any one of Claims 1 to 8, for screening antibiotics.



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20.Method for screening antibiotics, characterized in that it comprises at least the following steps:

a<sub>1</sub>)bringing a mini-PBP2x according to any one of Claims 1 to 7 into contact with a test substance,

b<sub>1</sub>)detecting, by any suitable means, the binding of said test molecule with the mini-PBP2x and/or the inhibition of the activity of said mini-PBP2x resulting from this binding, and

c<sub>1</sub>) selecting and identifying the active substances capable of binding to the mini-PBP2x and/or of inhibiting the activity of said mini-PBP2x, which can be used as antibiotics.

21.Method for identifying antibiotics, charac15 terized in that it comprises at least the following steps:

a<sub>2</sub>)preparing crystals from a mini-PBP2x according to any one of Claims 1 to 7,

b<sub>2</sub>) determining the three-dimensional structure

20 of said mini-PBP2x from the crystal obtained in a<sub>2</sub>), and

c<sub>2</sub>) identifying active substances capable of

binding to the mini-PBP2x and/or of inhibiting the

activity of said mini-PBP2x, which can be used as anti-

biotics.

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22. Screening kit for implementing the method according to Claim 20 or Claim 21, characterized in that it includes at least one protein, according to any one of Claims 1 to 8.

23. Screening kit for implementing the method according to Claim 20 or Claim 21, characterized in that it further includes at least one antibody according to claim 10.

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